

001080 55 200500

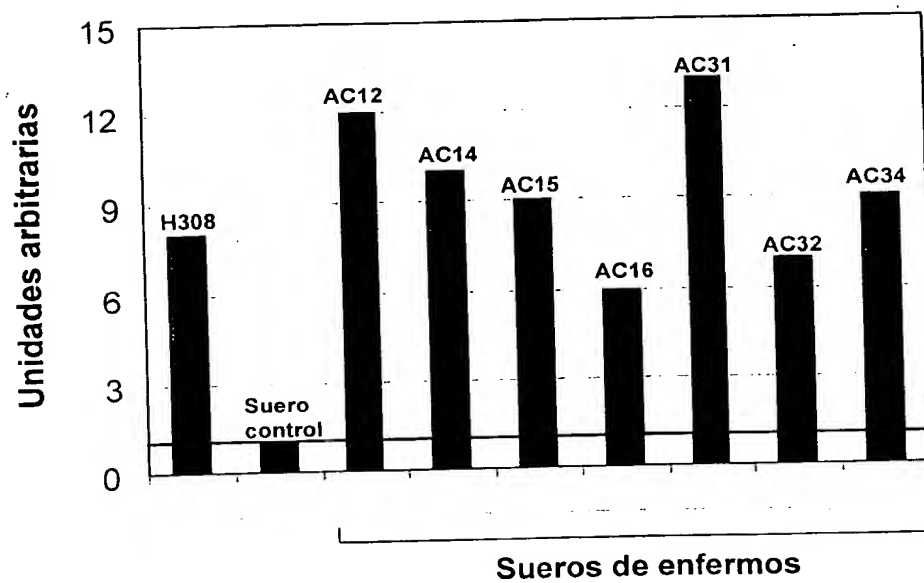
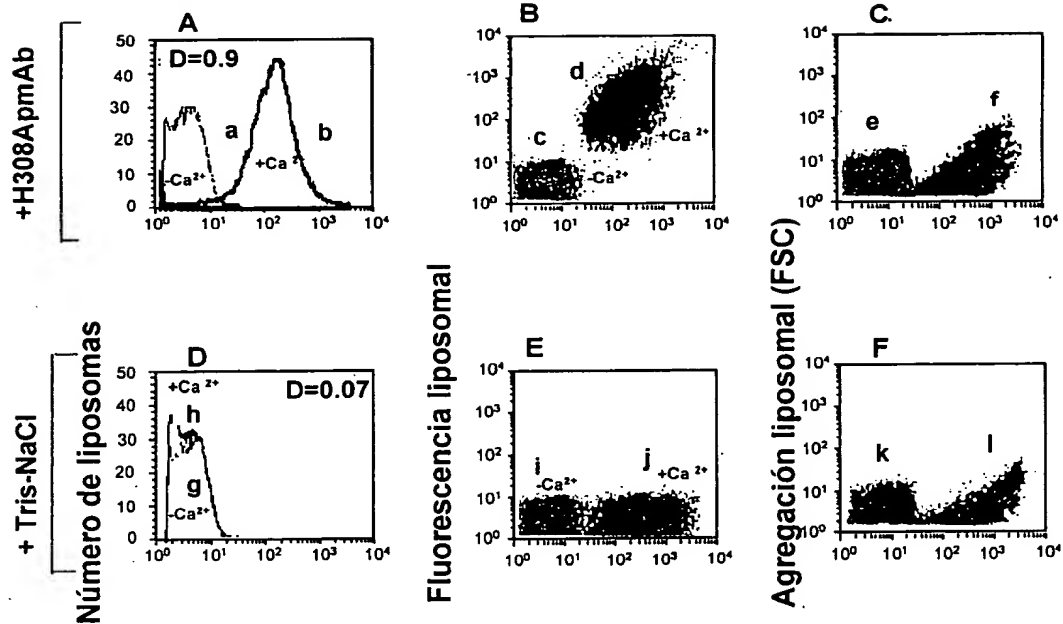


Figura 1

Liposomas de PC:PA



Liposomas de DPPC:PC:DPPA (1.2:0.8:1)

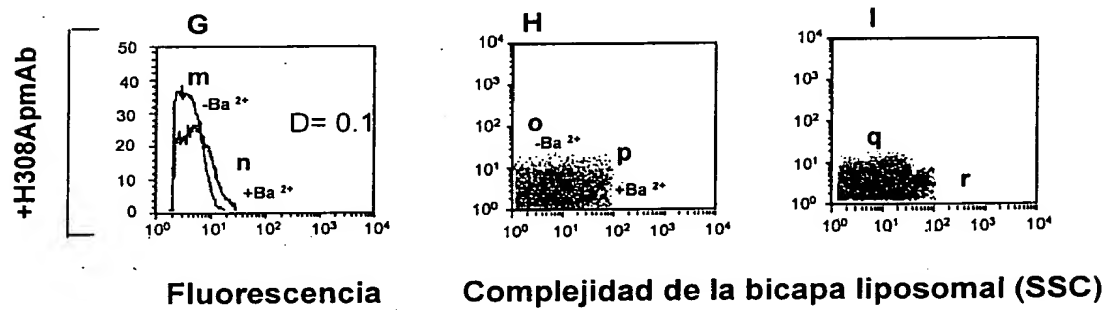


Figura 3

00000-9622960

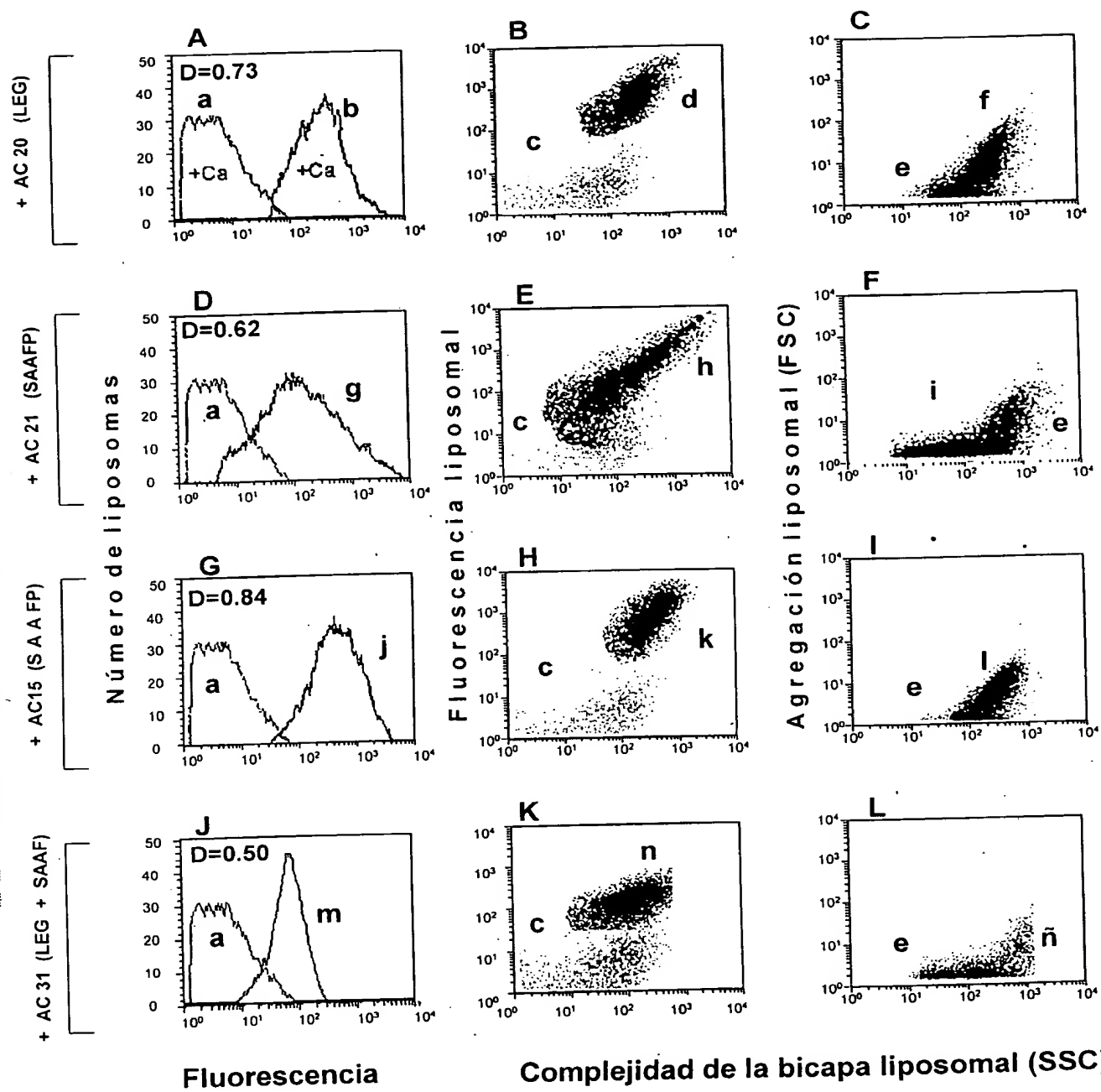
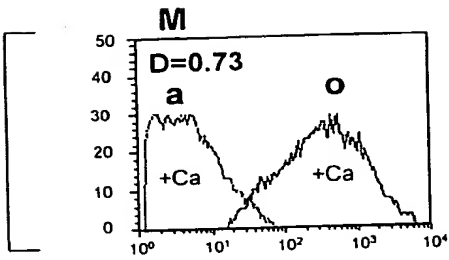


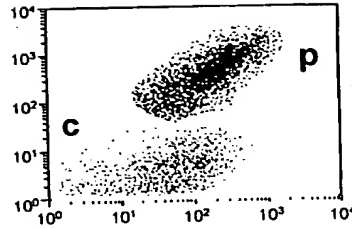
Figura 4 A

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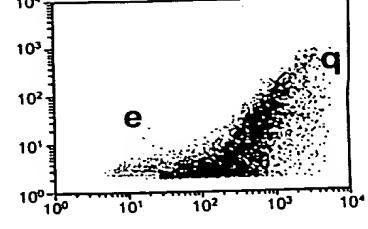
+ AC19 (LEG)



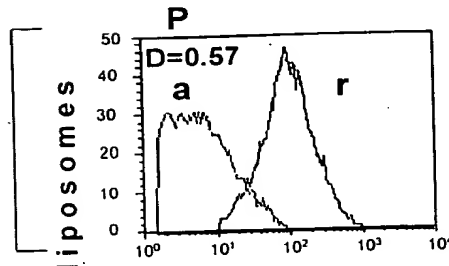
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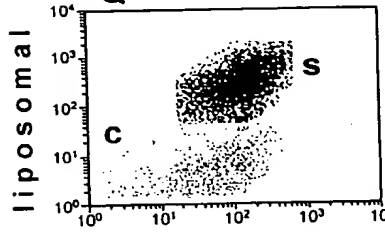
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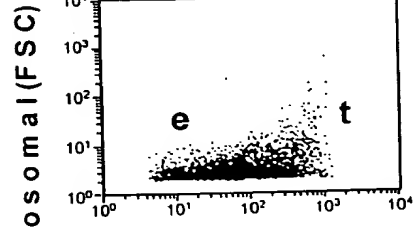
+ AC30 (SAAFP)



Q

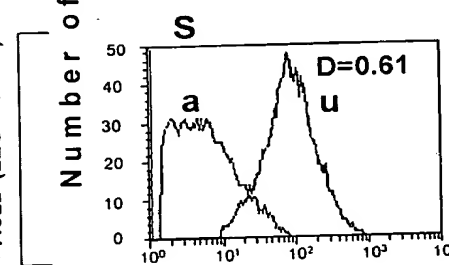


R

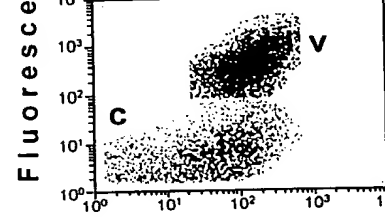


Number of liposomes

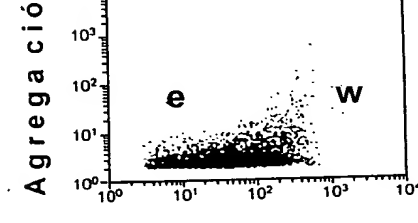
+ AC22 (LEG + SAAF)



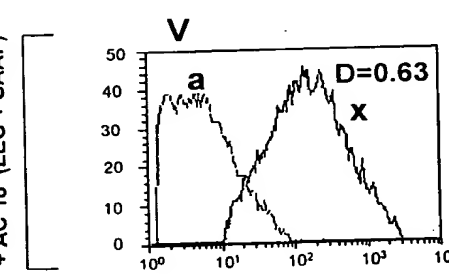
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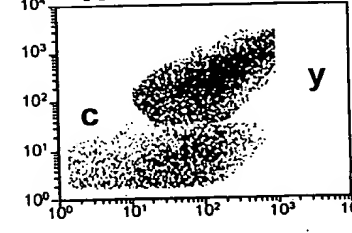
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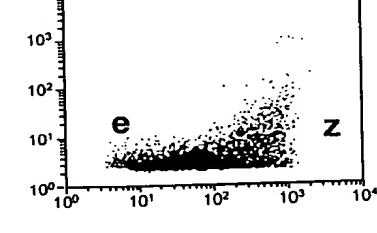
+ AC18 (LEG + SAAF)



W



X



Fluorescencia

Complejidad de la bicapa liposomal (SSC)

Figura 4 B

+AC15 (SAAFP)

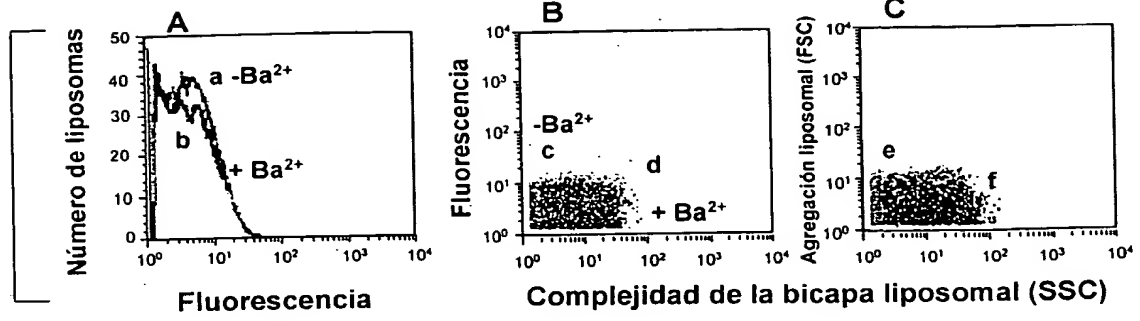


Figura 5

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001080-5466560

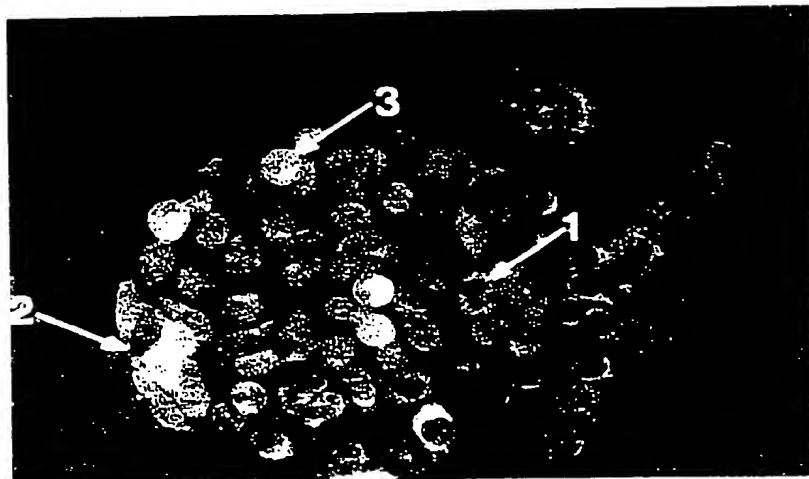


Figura 6

Sueros de enfermos	Sueros sin absorber (Unidades Arbitrias)	Sueros absorbidos con partículas lipídicas (Unidades Arbitrias)
Control	~1.3	~1.1
AC15	~3.1	~1.1
AC19	~3.7	~1.3
AC29	~1.6	~1.2
AC30	~1.6	~0.7
AC33	~4.1	~1.4
AC34	~2.2	~0.8

/

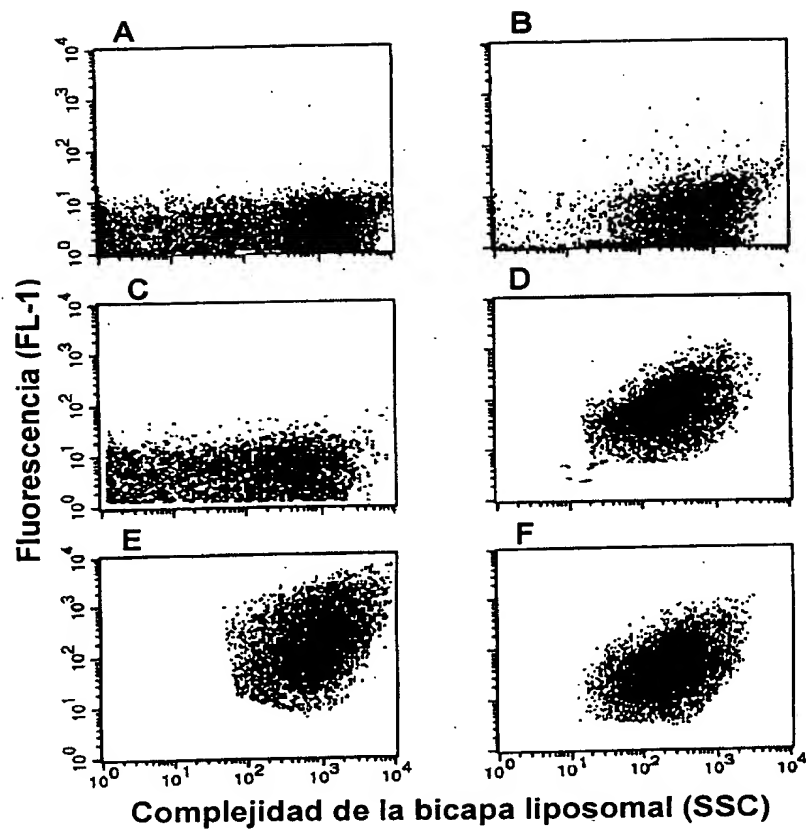


Figura 8

Figure 1 consists of six flow cytometry scatter plots arranged in a 3x2 grid, labeled A through F. Each plot has a logarithmic scale on both the x-axis and y-axis, ranging from 10^0 to 10^4 . The x-axis for all plots is labeled 'Complejidad de la bicapa liposomal (SSC)' and the y-axis is labeled 'Agregación liposomal (FSC)'. The plots show a positive correlation between the two variables, with the density of data points increasing as both values increase. Plot A shows a dense cluster of points at the bottom right, while plot F shows a more dispersed distribution of points across the plot area.

Figura 9

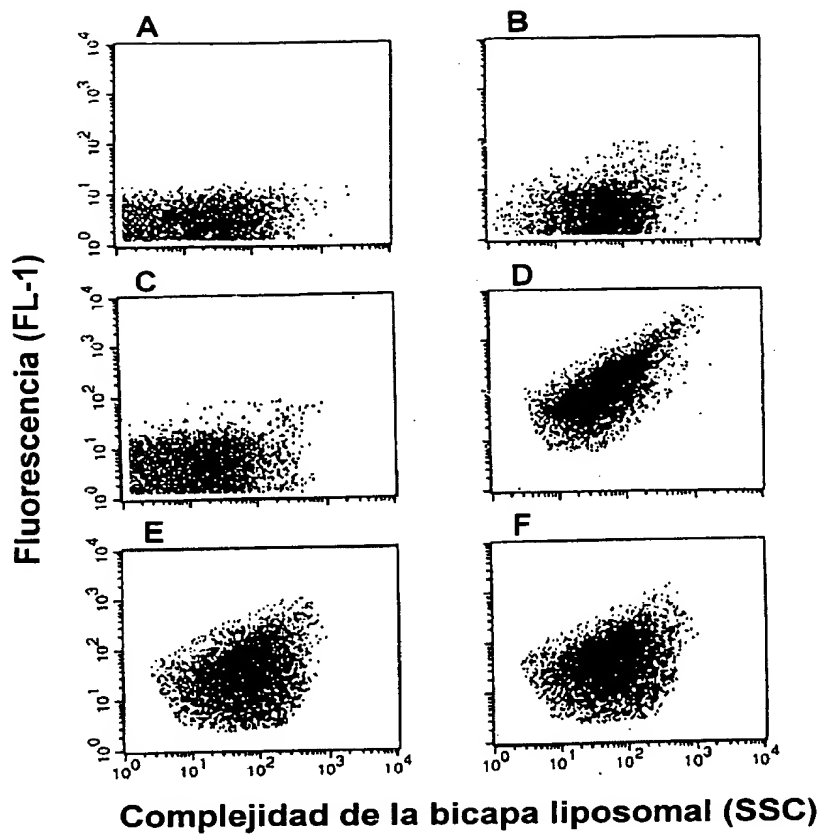


Figura 10

000000:52222250

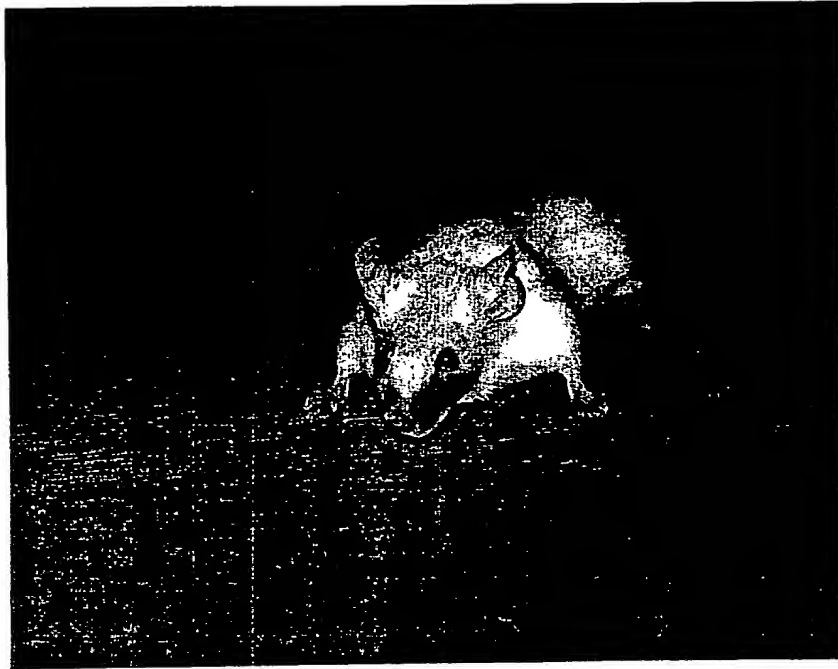


Figura 11

004060" 5522300

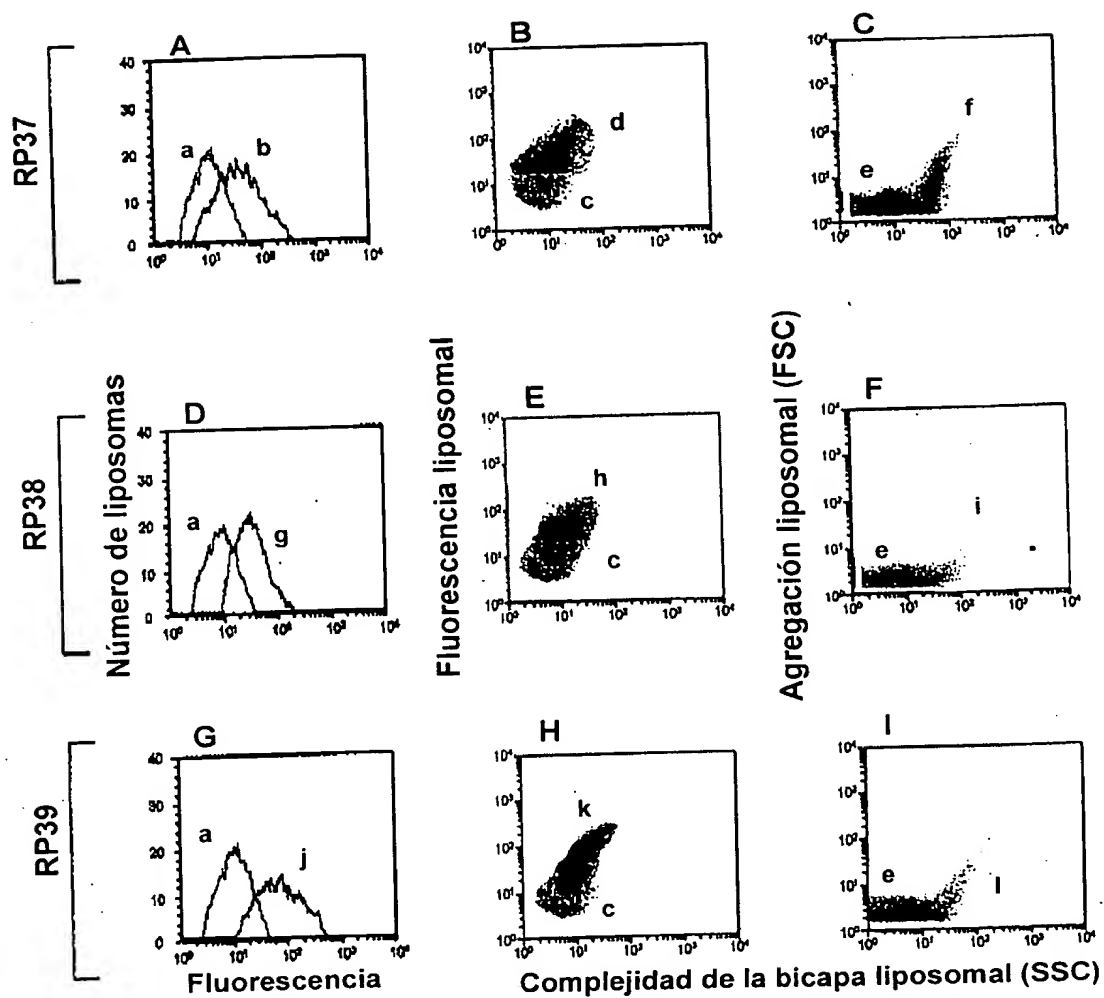
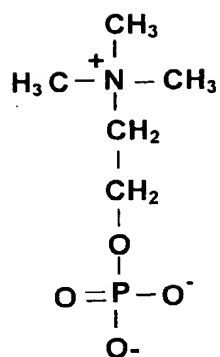
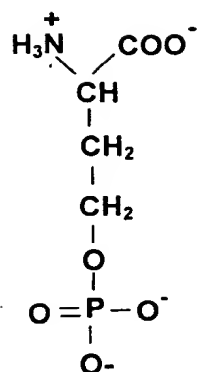


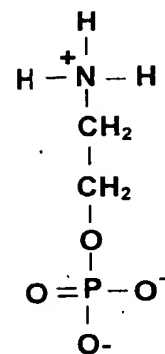
Figura 12



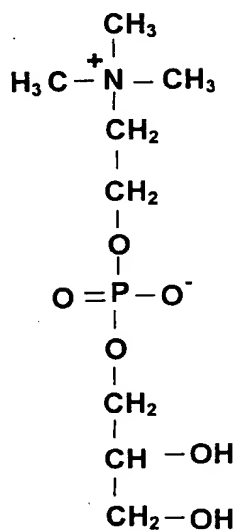
Fosforilcolina



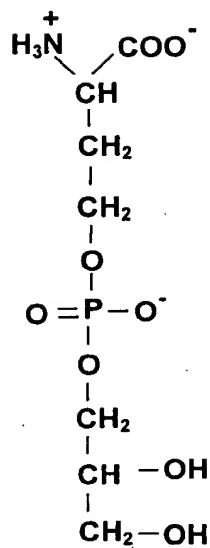
Fosforilserina



Fosforiletanolamina



Glicerolfosforilcolina



Glicerolfosforilserina

Figura 13

001030-5223300

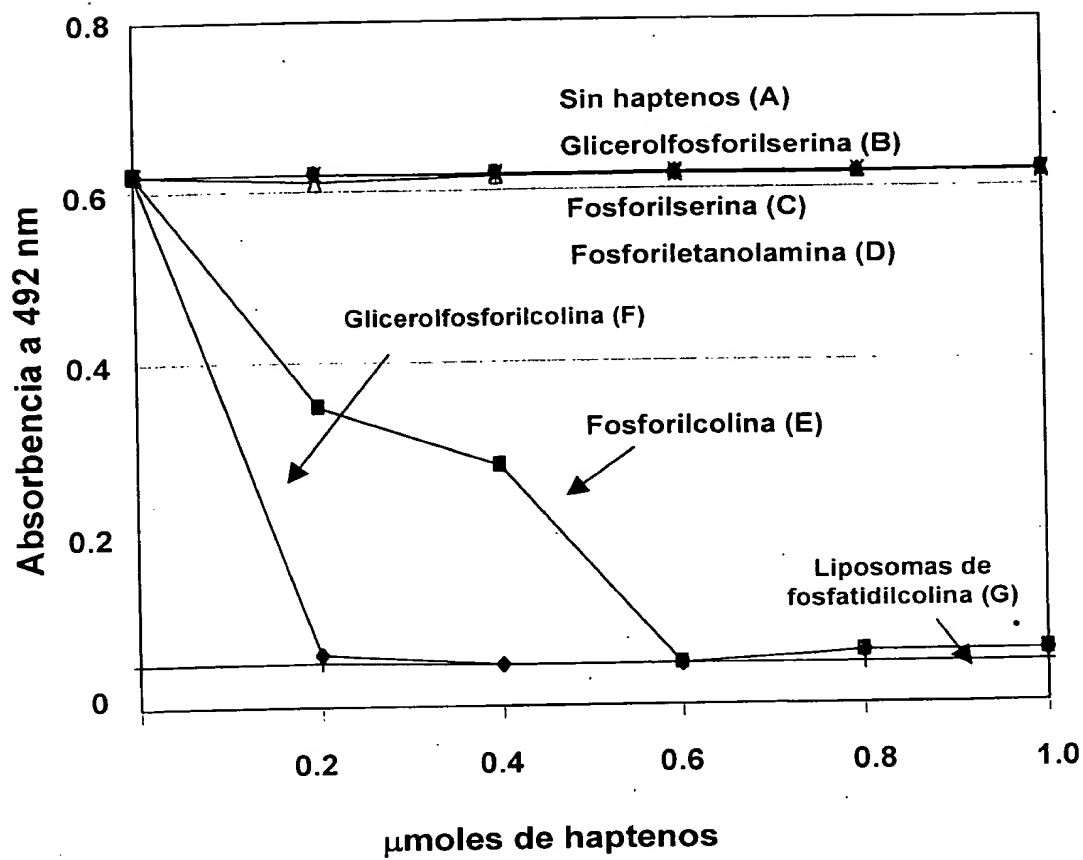


Figura 14

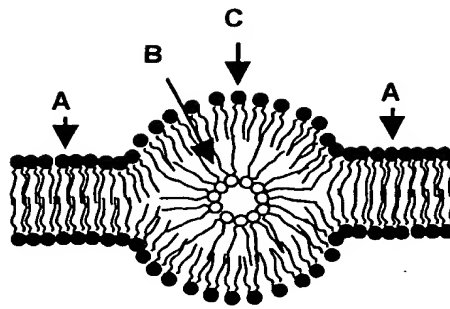


Figura 15

000000-000000

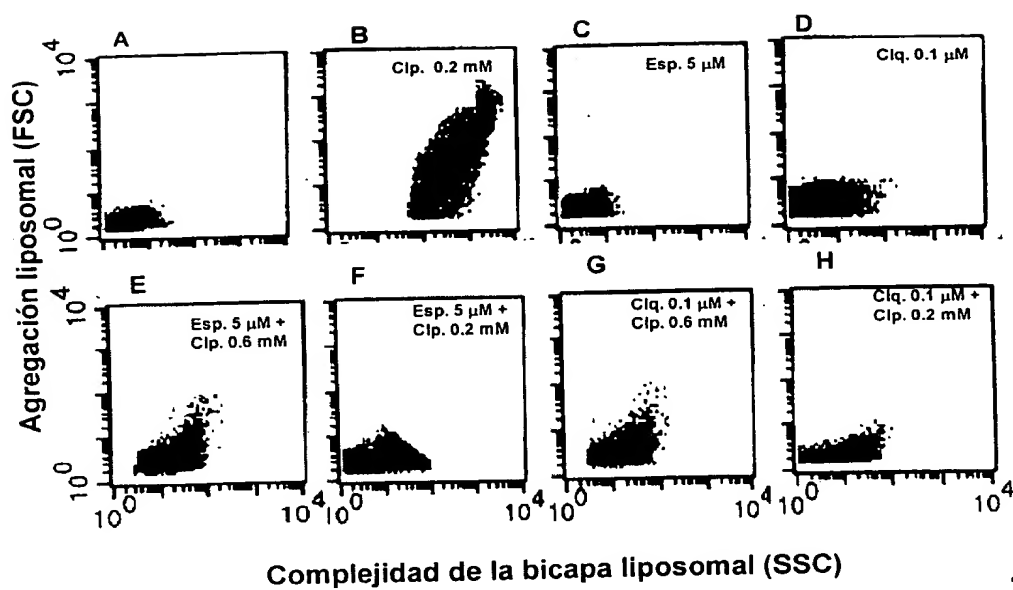


Figura 16